

Corrigenda: Genomic characterisation of *Arachis porphyrocalyx* (Valls & C.E. Simpson, 2005) (Leguminosae): multiple origin of *Arachis* species with $x = 9$. Comparative Cytogenetics 11(1): 29–43. doi: 10.3897/CompCytogen.v11i1.10339

Silvestri María Celeste¹, Alejandra Marcela Ortiz^{1,2}, Germán Ariel Robledo^{1,2}, José Francisco Montenegro Valls³, Graciela Inés Lavia^{1,2}

1 Instituto de Botánica del Nordeste (CONICET-UNNE, Fac. Cs. Agrarias), Sargento Cabral 2131, C.C. 209, 3400 Corrientes, Argentina **2** Facultad de Ciencias Exactas y Naturales y Agrimensura, UNNE, Av. Libertad 5460, 3400 Corrientes, Argentina **3** Embrapa Recursos Genéticos e Biotecnologia, Brasília, DF, Brasil

Corresponding author: Graciela Inés Lavia (graciela.lavia@yahoo.com.ar)

Academic editor: V.G. Kuznetsova | Received 10 October 2017 | Accepted 17 November 2017 | Published 8 December 2017

<http://zoobank.org/9109B40C-DB9D-433F-9113-41E553A9F6D9>

Citation: Silvestri MC, Ortiz AM, Robledo GA, Valls JFM, Lavia GI (2017) Corrigenda: Genomic characterisation of *Arachis porphyrocalyx* (Valls & C.E. Simpson, 2005) (Leguminosae): multiple origin of *Arachis* species with $x = 9$. Comparative Cytogenetics 11(1): 29–43. doi: 10.3897/CompCytogen.v11i1.10339. Comparative Cytogenetics 11(4): 819–820. <https://doi.org/10.3897/CompCytogen.v11i4.21560>

After the publication of our article, we detected an error in the figure 3. On the ideogram, the signal of 5S rDNA loci illustrated with a striped signal over the long arm of chromosome 2 at proximal/interstitial position is not observed. Correct figure is as follows:

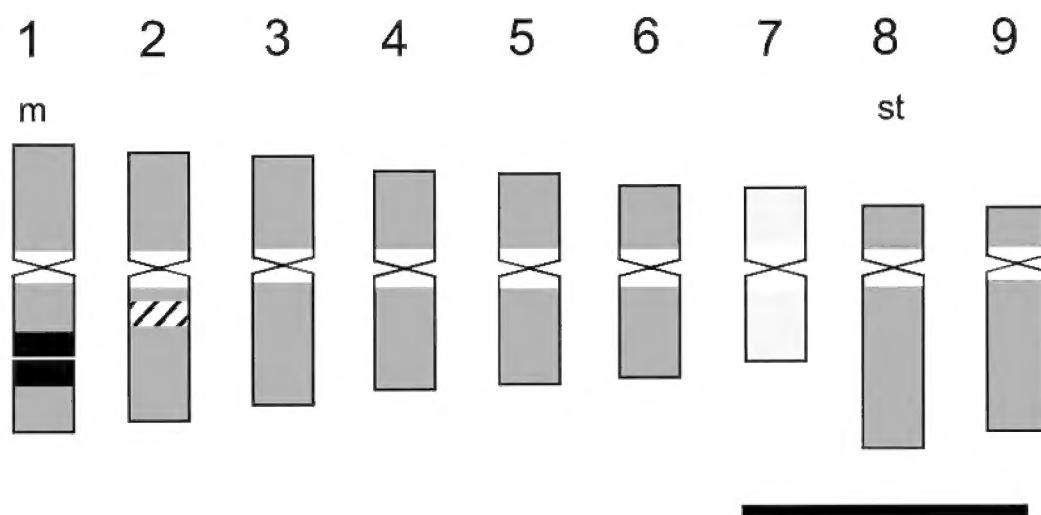


Figure 3. Ideogram of *A. porphyrocalyx* performed with measures of chromosomes obtained by classical technique. The “A” chromosome is represented with light grey colour. Distribution of 5S rDNA loci is illustrated with a striped signal and that of 18S–26S rDNA loci with a black signal. Heterochromatic regions counterstained with C-DAPI+ are represented with white bands. Scale = 2 μ m.